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Authors

Hogle, Shane L
Dupont, Christopher L
Hopkinson, Brian M
et al.

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Pervasive iron limitation at subsurface chlorophyll maxima of the California Current

Shane L. Hogle^{a,b,1}, Christopher L. Dupont^c, Brian M. Hopkinson^d, Andrew L. King^e, Kristen N. Buck^f, Kelly L. Roe^g, Rhona K. Stuart^h, Andrew E. Allen^{a,c}, Elizabeth L. Mannⁱ, Zackary I. Johnson^j, and Katherine A. Barbeau^{a,1}

^aScripps Institution of Oceanography, University of California, San Diego, La Jolla, CA 92093; ^bDepartment of Civil and Environmental Engineering, Massachusetts Institute of Technology, Cambridge, MA 02139; ^cMicrobial and Environmental Genomics, J. Craig Venter Institute, La Jolla, CA 92037; ^dDepartment of Marine Sciences, University of Georgia, Athens, GA 30602; ^eSection for Marine Biogeochemistry and Oceanography, Norwegian Institute for Water Research, NO-5006 Bergen, Norway; ^fCollege of Marine Science, University of South Florida, Tampa, FL 33620; ^gDepartment of Chemistry and Biochemistry, The College at Brockport, State University of New York, Brockport, NY 14420; ^hBiosciences and Biotechnology Division, Lawrence Livermore National Laboratory, Livermore, CA 94550; ⁱTrace Metal Biogeochemistry Laboratory, Bigelow Laboratory for Ocean Sciences, East Boothbay, ME 04544; and ^jMarine Laboratory and Biology Department, Duke University, Beaufort, NC, 28516

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Subsurface chlorophyll maximum layers (SCMLs) are nearly ubiquitous in stratified water columns and exist at horizontal scales ranging from the submesoscale to the extent of oligotrophic gyres. These layers of heightened chlorophyll and/or phytoplankton concentrations are generally thought to be a consequence of a balance between light energy from above and a limiting nutrient flux from below, typically nitrate (NO₃). Here we present multiple lines of evidence demonstrating that iron (Fe) limits or with light colimits phytoplankton communities in SCMLs along a primary productivity gradient from coastal to oligotrophic offshore waters in the southern California Current ecosystem. SCML phytoplankton responded markedly to added Fe or Fe/light in experimental incubations and transcripts of diatom and picoeukaryote Fe stress genes were strikingly abundant in SCML metatranscriptomes. Using a biogeochemical proxy with data from a 40-y time series, we find that diatoms growing in California Current SCMLs are persistently Fe deficient during the spring and summer growing season. We also find that the spatial extent of Fe deficiency within California Current SCMLs has significantly increased over the last 25 y in line with a regional climate index. Finally, we show that diatom Fe deficiency may be common in the subsurface of major upwelling zones worldwide. Our results have important implications for our understanding of the biogeochemical consequences of marine SCML formation and maintenance.

nutrient limitation | iron | light | California Current | deep chlorophyll maximum

Fe and light are essential for phytoplankton photosynthesis, but both resources are scarce in much of the ocean. Surface ocean primary productivity is limited by the availability of Fe in some regions (1), and mesoscale Fe fertilization experiments now firmly demonstrate that Fe availability controls phytoplankton biomass and growth rates in the Southern, equatorial Pacific, and subarctic Pacific oceans (2). In addition, phytoplankton Fe limitation has been observed in midlatitude coastal upwelling zones (3, 4), throughout mesoscale circulation features (5, 6), and at the edge of subtropical gyres (7). In the surface ocean light attenuates rapidly to less than 1% of incident photosynthetically available radiation (z_{1%}) at depths from 50 m to 200 m, depending on turbidity. However, many diverse phytoplankton groups have adapted to growth at depths approaching z_{1%} despite the challenging low-light conditions. Prior studies noting the overlapping scarcity of Fe and light in much of the ocean predicted that these two resources synergistically colimit phytoplankton growth (8), particularly in subsurface chlorophyll maximum layers (SCMLs) (9). Indeed, work with cultured phytoplankton demonstrates that Fe/light colimitation can arise when demand for Fe-rich photosynthetic redox proteins increases under low-light conditions (9, 10). However, the potential for phytoplankton Fe or Fe/light (co)limitation in SCMLs has been explored only in a handful of field studies despite the significant feedbacks linking Fe/light (co)limitation, dust deposition, and oceanic CO₂ uptake in global biogeochemical models

(11). Although Fe/light colimitation has been observed in some high-latitude SCMLs (12, 13), mid-/low-latitude SCMLs from both coastal and pelagic zones remain understudied. Dissolved Fe minima at SCMLs from the subtropical North Pacific gyre (14) and the Sargasso Sea (15) may be a consequence of intense biological demand, even Fe limitation, during summer months. One study documented phytoplankton Fe/light colimitation from mesotrophic and oligotrophic SCMLs in the California Bight and the eastern tropical North Pacific (16), while another found SCMLs in the oligotrophic Western Pacific to be mostly light limited with some groups of microbial eukaryotes potentially

Significance

The vertical distribution of phytoplankton cells and chlorophyll concentrations throughout the sunlit water column is rarely uniform. In many ocean regions, chlorophyll concentrations peak in distinct and persistent layers deep below the surface called subsurface chlorophyll maximum layers (SCMLs). SCML formation is hypothesized to reflect the consequences of phytoplankton light/macronutrient colimitation, behavior, and/or photoacclimation. We discovered unexpectedly persistent and widespread phytoplankton iron limitation and iron/light colimitation in SCMLs of the California Current and at the edge of the North Pacific Subtropical Gyre using shipboard incubations, metatranscriptomics, and biogeochemical proxies. These results suggest that interactions and feedbacks between iron and light availability play an important and previously unrecognized role in controlling the productivity and biogeochemical dynamics of SCMLs.

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Data deposition: All data supporting the findings of this study are available at the following websites: CalCOFI time series data have been deposited in the CalCOFI data archives, new.data.calcofi.org/index.php/reporteddata. Metatranscriptome and metagenome biological sequence files have been deposited in iMicrobe, <https://imicrobe.us> (iMicrobe project ID: CAM.P.0001069). Biogeochemical data subsets, metatranscriptome annotations, and all computer code required to reproduce the results reported in this study have been deposited in Zenodo, <https://doi.org/10.5281/zenodo.1495558> and <https://doi.org/10.5281/zenodo.1495504>. Unprocessed biogeochemical data have been deposited in the UCSD DataZoo Research Project, oceaninformatics.ucsd.edu/datazoo/catalogs/ccelcer/sources/1758.

¹To whom correspondence may be addressed. Email: shogle@mit.edu or kbarbeau@ucsd.edu.

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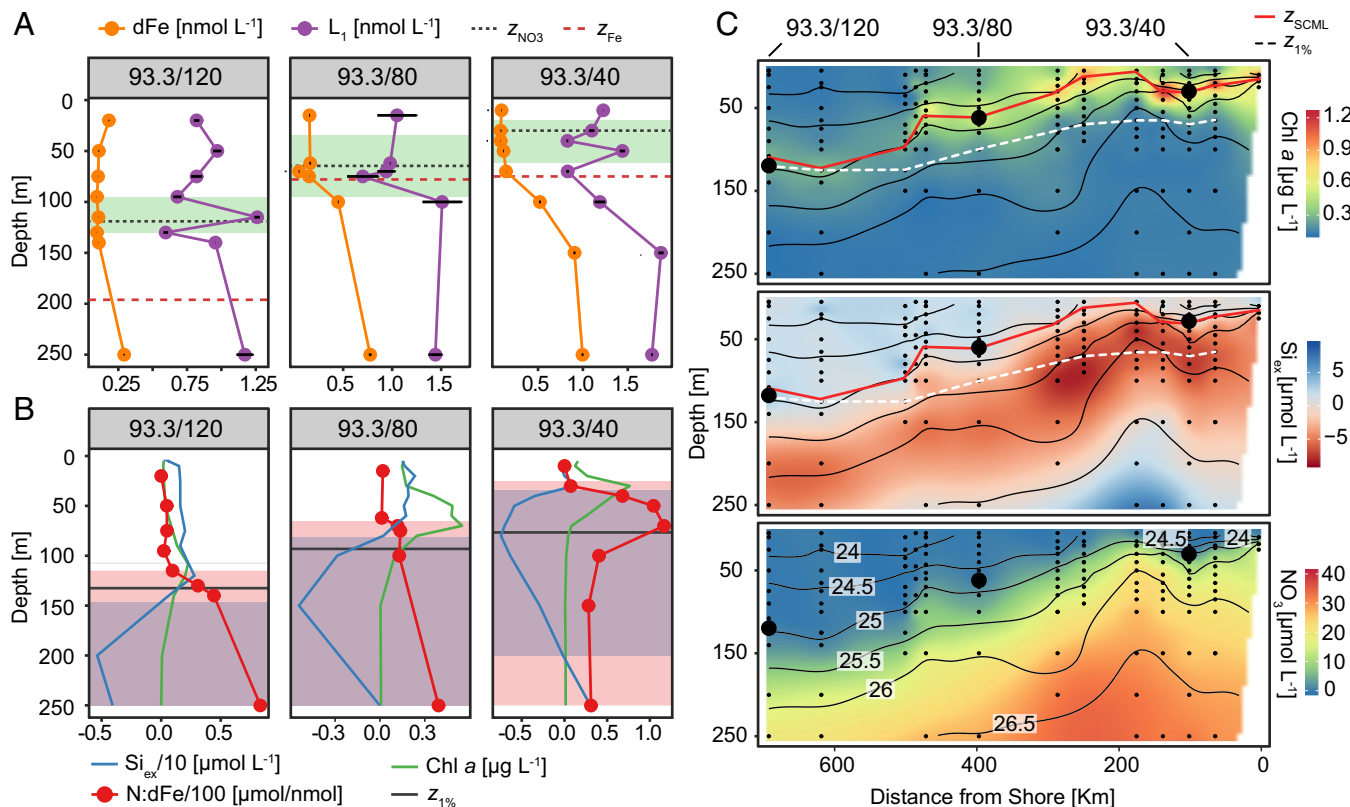


Fig. 1. (A) Profiles of dissolved Fe (orange) and L_1 (purple) concentrations. The green box denotes depths where chlorophyll *a* (Chl *a*) is within 50% of the concentration at the SCML depth (z_{SCML}), the dashed black line is nitrate line depth (z_{NO_3}), and the dashed red line is ferricline depth (z_{Fe}) (SI Appendix, Table S1 and section 4). (B) Profiles of Chl *a* (green), $Si_{ex}/10$ (blue), and $N:Fe/100$ (red). The black line represents the depth of 1% of incident irradiance ($z_{1\%}$). Red ($N:Fe \geq 8$), blue ($Si_{ex} < 0$), and purple (both proxies) indicate depth ranges with potential diatom Fe deficiency. (C) Chl *a*, NO_3 , and Si_{ex} ($26.5 \text{ kg} \cdot \text{m}^{-3}$ source isopycnal) sections. Potential density anomaly (σ_θ) contours are in black. Small circles show nutrient sampling depths and large circles show incubation and metatranscriptome depths. White and red dashed lines show $z_{1\%}$ and z_{SCML} , respectively.

over control), and total Chl *a* (5.5-fold over control) increases, while responses to Fe or light alone were similar to each other and significantly smaller than for Fe + light. However, the single addition of iron or light significantly enhanced NO₃ drawdown (Fe, 1.7-fold over control; light, 1.5-fold over control) and the addition of iron significantly increased the Chl *a* concentration (2.5-fold over control). Large chain-forming diatoms dominated incubation responses to added Fe + light (*SI Appendix, Fig. S4*), while iron-stress transcripts were in the top 1% of all expressed *Phaeocystis* and pelagophyte genes in situ (Fig. 3). Incubation experiments from 93.3/120 displayed a roughly twofold increase in diatom cell numbers, NO₃ drawdown, Chl *a* concentrations, and primary production in Fe + light conditions over the control (Fig. 2). Only the increase in Chl *a* was statistically significant, but it was challenging to detect significant differences in the means across incubation replicates because responses in the offshore zone were substantially smaller and noisier than in other regions. Given this caveat, our results suggest a potential for simultaneous Fe/light colimitation in oligotrophic SCMLs in the offshore zone. Smaller non-chain-forming diatoms were the dominant responders to the offshore zone Fe addition incubations (*SI Appendix, Fig. S4*), and the in situ positive Si_{ex} signal at the offshore SCML suggests that heavily silicified, Fe-limited forming diatoms were not abundant at the time of sampling. Indeed, pelagophytes numerically dominated the metatranscriptomes at 93.3/120, and pelagophyte pTf/ISIP3 transcripts were in the top 1% of all expressed transcripts in the metatranscriptome and in the top 0.1% of pelagophyte transcripts indicating significant cellular resource investment into Fe acquisition by small, nonsilicifying phytoplankton.

SCML Fe Deficiency Estimated from a 40-y Time Series. We sought to characterize the potential for diatom Fe deficiency in the southern CC across broader spatial and temporal scales by leveraging 40 y of monthly sampling data collected from 75 stations distributed over a 190,000-km² area as a part of the CalCOFI program (calcofi.org). We used Si_{ex} as a biogeochemical proxy for diatom Fe deficiency because H₄SiO₄ and NO₃ measurements are readily available in the CalCOFI dataset and there is a strong correlation between negative Si_{ex} and experimentally determined Fe limitation in our results (Figs. 1 and 2 and *SI Appendix, Fig. S3*) and the results of others (5, 6, 24). The Si_{ex} tracer assumes minimal effects from horizontal mixing/advection, shifts in upwelling source depth, and other processes (nitrification, denitrification, variability in Si:N remineralization ratios) that may integrate non-specific biogeochemical signals. In most settings, such as in the euphotic zone of the CC, these assumptions appear to be valid, but we note that in other biogeochemical regimes they may not be. In all test cases and datasets we examined Si_{ex} performed as a robust tracer (*SI Appendix, section 13A*).

During the spring and summer months over the last 40 y at least 30% of all SCML samples in the southern CC were Fe deficient to some degree ($\text{Si}_{\text{ex}} < 0$). Fe deficiency is disproportionately concentrated in SCMLs from the inshore (43% negative) and transition zones (26% negative) compared with the offshore zone (7% negative) (Fig. 4 and [SI Appendix, Fig. S6](#)). On average Si_{ex} from inshore and transition zone SCMLs has steadily become more negative since 2000 in contrast to the general increase in the 1990s and most of the 1980s ([SI Appendix, Fig. S7](#)). We also find that the total spatial area of Fe-deficient SCMLs has significantly increased for the inshore



Figure 1 displays taxonomically resolved RPKM values for 16S rRNA sequences across four taxonomic groups (Diatoms, Dinoflagellates, Pelagophytes, and Phaeocystis) and three different 16S rRNA sequences (93.3/120, 93.3/80, and 93.3/40). The y-axis represents RPKM values on a logarithmic scale from 10^1 to 10^5 . The x-axis lists the genes: pTf/ISIP3, Flavodoxin, PR-like, NRT1, HSP, RIBO, RUBISCO, and PS I/II rel. A dashed horizontal line at 10^2 indicates the detection limit. The data shows that RPKM values are generally higher for the 93.3/120 sequence compared to the 93.3/80 and 93.3/40 sequences, particularly for the Diatoms and Dinoflagellates groups.

Fig. 3. Relative abundance of transcripts/isoforms from known iron-stress genes and genes related to essential cellular processes. The vertical axis (log10 scale) represents the reads per kilobase of transcript per million mapped reads to each of the four taxonomic groups at each station. Individual assembled transcripts/isoforms are binned into functional groups (horizontal axis). Functional groups with greater than 10 unique transcripts/isoforms are represented by boxplots; otherwise expression values for individual transcripts/isoforms are displayed. Individual transcripts/isoforms above the solid black lines exceed the taxonomic and library-specific 95th percentile ranking while those above the dashed line exceed the 99th percentile ranking and those in the lower 10% of percentile ranking are not included. HSP, heat-shock proteins; NRT1, nitrate transporter; PR-like, eukaryotic proteorhodopsin-like; PS I/II rel., photosystem reaction center proteins (PsbE, -F, -H-N, -R, -S, -W) and light-harvesting complex proteins (Chl *a/b/c* or fucoxanthin binding); RIBO, all ribosomal assembly protein subunits; RUBISCO, ribulose-1,5-bisphosphate carboxylase.

PNAS | December 26, 2018 | vol. 115 | no. 52 | 13303

CC under anthropogenic climate change (48, 49), which may drive diatom communities at the SCML toward Fe limitation if the associated Fe fluxes do not increase proportionally. This potential atmospheric–biogeochemical linkage demonstrates a connection, mediated by iron, by which the changing climate may influence carbon cycling and primary productivity in SCMLs of the CC and potentially other eastern boundary currents.

Materials and Methods

Method details are available in *SI Appendix*. Samples for total dissolved iron concentrations, iron-binding ligand concentrations, and incubation experiments were collected and processed as described in refs. 4 and 24. Dissolved Fe was measured as in ref. 4 and Fe-binding ligands were measured as in ref. 50. The transcriptomic data (including all relevant methods) were introduced in a prior publication (26). Macronutrients were sampled using a rosette sampler and analyzed following the standard operating procedures from the California Current Ecosystem Long-Term Ecological Research program. Triplicate or duplicate 4-L incubations were conducted in acid-cleaned polycarbonate bottles, housed in a Percival incubator at 16 °C with a 12:12 light:dark cycle at a high- and low-light treatment and added Fe (*SI Appendix, Table S3*). Primary productivity and photo-

physiology were measured as described in refs. 17 and 26. CalCOFI and *World Ocean Atlas* hydrographic and nutrient data were downloaded from new.data.calcofi.org and <https://www.nodc.noaa.gov>. The Si_{ex} proxy (*SI Appendix, section 13A*) at SCML depths was calculated as $Si_{ex} = [\mu\text{mol H}_4\text{SiO}_4 \cdot \text{L}^{-1}] - ([\mu\text{mol NO}_3 \cdot \text{L}^{-1}] \times R_{Si:NO_3})$, where $R_{Si:NO_3}$ is the micromolar ratio of H_4SiO_4 to NO_3 at $\sigma_\theta = 25.8 \text{ kg} \cdot \text{m}^{-3}$ or $26.5 \text{ kg} \cdot \text{m}^{-3}$.

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